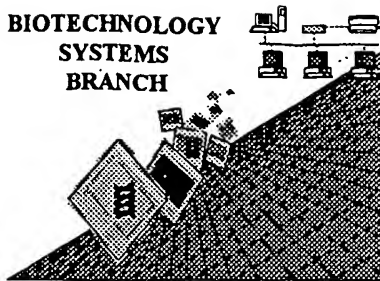


227
0460

BIOTECHNOLOGY
SYSTEMS
BRANCH



RAW SEQUENCE LISTING
ERROR REPORT

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number: 10/033,399
Source: 01PE
Date Processed by STIC: 1/18/2002

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FOR CRF SUBMISSION QUESTIONS, PLEASE CONTACT MARK SPENCER, 703-308-4212.

FOR SEQUENCE RULES INTERPRETATION, PLEASE CONTACT ROBERT WAX, 703-308-4216.

PATENTIN 2.1 e-mail help: patin21help@uspto.gov or phone 703-306-4119 (R. Wax)

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TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE **CHECKER**
VERSION 3.1 PROGRAM, ACCESSIBLE THROUGH THE U.S. PATENT AND
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<http://www.uspto.gov/web/offices/pac/checker>

Applicants submitting genetic sequence information electronically on diskette or CD-Rom should be aware that there is a possibility that the disk/CD-Rom may have been affected by treatment given to all incoming mail.

Please consider using alternate methods of submission for the disk/CD-Rom or replacement disk/CD-Rom, including:

1. EFS-Bio (<<http://www.uspto.gov/ebs/efs/downloads/documents.htm>> , EFS Submission

User Manual - ePAVE)

2. U.S. Patent and Trademark Office, Box Sequence, P.O. Box 2327, Arlington, VA 22202

3. Hand Carry directly to:

U.S. Patent and Trademark Office, Technology Center 1600, Reception Area, 7th Floor, Examiner Name,
Sequence Information, Crystal Mall One, 1911 South Clark Street, Arlington, VA 22202

Or

U.S. Patent and Trademark Office, Box Sequence, Customer Window, Lobby, Room 1B03, Crystal Plaza Two,
2011 South Clark Place, Arlington, VA 22202

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Box Sequence, Room 1B03-Mailroom, Crystal Plaza Two, 2011 South Clark Place, Arlington, VA 22202



OIPE

RAW SEQUENCE LISTING

PATENT APPLICATION: US/10/033,399

DATE: 01/18/2002

TIME: 08:04:15

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Output Set: N:\CRF3\01182002\J033399.raw

PP-5-4

Does Not Comply
Corrected Diskette Needed

C-11
C-11

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3 <110> APPLICANT: Wang, Caili
4     Zhong, Pingyu
5     Wang, Xinwei
7 <120> TITLE OF INVENTION: ADAPTER-DIRECTED DISPLAY SYSTEMS
9 <130> FILE REFERENCE: 13403.0005NPUS00
11 <140> CURRENT APPLICATION NUMBER: US/10/033,399
11 <141> CURRENT FILING DATE: 2001-11-02
11 <160> NUMBER OF SEQ ID NOS: 24
13 <170> SOFTWARE: PatentIn version 3.1
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16 <211> LENGTH: 57
17 <212> TYPE: DNA
18 <213> ORGANISM: Bacteriophage M13
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51 <213> ORGANISM: Artificial Sequence
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54 <223> OTHER INFORMATION: Synthetic, comprising phage gene III leader sequence, GABAB
recep
55     tor 2 domain and Myc domain
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62 cagctgcagg acgtcggagg ttgcgcggcc gcagaacaaa aactcatctc agaagaggat      180
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68 <211> LENGTH: 74

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RAW SEQUENCE LISTING

DATE: 01/18/2002

PATENT APPLICATION: US/10/033,399

TIME: 08:04:15

Input Set : A:\13403.0005NPUS00.ST25.txt

Output Set: N:\CRF3\01182002\J033399.raw

69 <212> TYPE: PRT

70 <213> ORGANISM: Artificial Sequence

72 <220> FEATURE:

73 <223> OTHER INFORMATION: Synthetic, comprising phage gene III leader sequence, GABAB
recep

74 tor 2 domain and Myc domain

76 <400> SEQUENCE: 5

78 Leu Val Val Pro Phe Tyr Ser His Ser Ala Thr Ser Arg Leu Glu Gly

79 1 5 10 15

82 Leu Gln Ser Glu Asn His Arg Leu Arg Met Lys Ile Thr Glu Leu Asp

83 20 25 30

86 Lys Asp Leu Glu Glu Val Thr Met Gln Leu Gln Asp Val Gly Gly Cys

87 35 40 45

90 Ala Ala Ala Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu Arg Ser Gly

91 50 55 60

94 Gly Gly Thr Val Glu Ser Cys Leu Ala Lys

95 65 70

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99 <211> LENGTH: 56

100 <212> TYPE: PRT

101 <213> ORGANISM: Artificial Sequence

103 <220> FEATURE:

104 <223> OTHER INFORMATION: Synthetic, comprising phage gene III leader sequence, GABAB
recep

105 tor 2 domain and Myc domain

107 <400> SEQUENCE: 6

109 Thr Ser Arg Leu Glu Gly Leu Gln Ser Glu Asn His Arg Leu Arg Met

110 1 5 10 15

113 Lys Ile Thr Glu Leu Asp Lys Asp Leu Glu Glu Val Thr Met Gln Leu

114 20 25 30

117 Gln Asp Val Gly Gly Cys Ala Ala Glu Gln Lys Leu Ile Ser Glu

118 35 40 45

121 Glu Asp Leu Arg Ser Gly Gly Gly

122 50 55

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127 <212> TYPE: DNA

128 <213> ORGANISM: Artificial Sequence

130 <220> FEATURE:

131 <223> OTHER INFORMATION: Synthetic, comprising ampicillin gene sequence, ColeI
replication

132 origin, fl replication origin, Plac promoter, GABAB receptor 1 d

133 omain, histidine tag

135 <400> SEQUENCE: 7

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138 gcttccggct cgtatgttgt gtggaattgt gagcgataa caatttaccg gttctttaag 120

140 gaggaattaa aaaatgaaat acctattgcc tacggcagcc gctggattgt tattactcgc 180

142 ggcccagccg gccatggcgg ccctgcaggc ctctagagcg gccgctggag gtgaggagaa 240

144 gtcccggctg ttggagaagg agaaccgtga actggaaaag atcattgctg agaaagagga 300

146 gcgtgtctct gaactgcgcc atcaactcca gtctgttagga ggtttagat cttatccata 360

148 cgacgtacca gactacgcag gaggtcatca ccattcatcac cattaatgag tcgacctcga 420

150 ccaattcgcc ctatagttag tcgtattaca attcactggc cgtcgtttta caacgtcgtg 480

RAW SEQUENCE LISTING

DATE: 01/18/2002

PATENT APPLICATION: US/10/033,399

TIME: 08:04:15

Input Set : A:\13403.0005NPUS00.ST25.txt

Output Set: N:\CRF3\01182002\J033399.raw

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156 atggcgaatg ggacgcgccc tgtagcggcg cattaagcgc ggcgggtgtg gtggttacgc 660
158 gcagcgtgac cgctacactt gccagcgcgc tagcgccgcg tcctttcgtt ttcttccctt 720
160 cttttctcgc cacgttcgcc ggctttcccc gtcaagctct aaatcggggg ctccctttag 780
162 ggttccgatt tagtgcttta cggcacctcg accccaaaaa acttgattag ggtgatggtt 840
164 cacgtagtgg gccatcgccc tgatagacgg tttttcgccc tttgacgttg gagtccacgt 900
166 tctttaatag tggactcttg ttccaaactg gaacaacact caaccctatc tcggtctatt 960
168 cttttgattt ataagggatt ttgccgattt cggcctattg gttaaaaaat gagctgattt 1020
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172 tcgggggaaat gtgcgcggaa cccctatttg tttatttttc taaatacatt caaatatgta 1140
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176 gagtattcaa catttccgtg tcgcccttat tccctttttt gcggcatttt gccttccgtt 1260
178 ttttgctcac ccagaaacgc tggtgaaagt aaaagatgct gaagatcagt tgggtgcacg 1320
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182 agaacgtttt ccaatgatga gcaactttta agttctgcta tgtggcgcgg tattatcccg 1440
184 tattgacgcc gggcaagagc aactcggtcg ccgcatacac tattctcaga atgacttggt 1500
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196 ccggcaacaa ttaatagact ggatggaggg ggataaagtt gcaggaccac ttctgcgctc 1860
198 ggcccttccg gctggctggt ttattgctga taaatctgga gccggtgagc gtgggtctcg 1920
200 cggtatcatt gcagcactgg ggccagatgg taagccctcc cgtatcgtag ttatctacac 1980
202 gacggggagt caggcaacta tggatgaacg aaatagacag atcgtctgaga taggtgcctc 2040
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206 aaaacttcat ttttaattta aaaggatcta ggtgaagatc ctttttgata atctcatgac 2160
208 caaaatccct taacgtgagt ttctgttcca ctgagcgtca gaccccgtag aaaagatcaa 2220
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214 aactggcttc agcagagcgc agataccaaa tactgtcctt ctagtgtagc cgtagttagg 2400
216 ccaccacttc aagaactctg tagcaccgcc tacatacctc gctctgctaa tcctgttacc 2460
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234 taccgctcgc cgcagccgaa cgaccgagcg cagcgagtca gtgagcgagg aagcggaaga 3000
236 gcgccaata cgcaaaccgc ctctccccgc gcgttggcgg attcattaat gcagctggca 3060
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242 <211> LENGTH: 192
243 <212> TYPE: DNA
244 <213> ORGANISM: Bacteriophage M13
246 <400> SEQUENCE: 8

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RAW SEQUENCE LISTING

PATENT APPLICATION: US/10/033,399

DATE: 01/18/2002

TIME: 08:04:15

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Output Set: N:\CRF3\01182002\J033399.raw

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 249 aaactcatct cagaagagga tctgagatct agatctggag gcggtactgt tgaaagttgt 120
 251 ttagcaaaac ctcatacaga aaattcattt actaacgtct ggaaagacga caaaacttta 180
 253 gatcgttacg ct 192

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257 <211> LENGTH: 64

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261 <220> FEATURE:

262 <221> NAME/KEY: MISC_FEATURE

263 <222> LOCATION: (11)..(11)

264 <223> OTHER INFORMATION: Xaa = stop codon

267 <400> SEQUENCE: 9

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 270 1 5 10 15
 273 Ala Ala Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu Arg Ser Arg Ser
 274 20 25 30
 277 Gly Gly Gly Thr Val Glu Ser Cys Leu Ala Lys Pro His Thr Glu Asn
 278 35 40 45
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 282 50 55 60

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286 <211> LENGTH: 2962

287 <212> TYPE: DNA

288 <213> ORGANISM: Artificial Sequence

290 <220> FEATURE:

291 <223> OTHER INFORMATION: Synthetic, comprising ampicillin gene sequence, ColE1
 replication

292 origin, fl replication origin, Plac promoter, influenza virus he

293 magglutinin tag

295 <400> SEQUENCE: 10

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 300 ctttagtaag gaggaattaa aaaatgaaat acctattgcc tacggcagcc gctggattgt 180
 302 tattactcgc ggcccagccg gccatggcgg cctgcaggc ctctagagcg gccgcttacc 240
 304 cgtacgacgt tccggactac gcaggtggct gctgataagt cgacctcgac caattcgccc 300
 306 tatagtgaat cgtattacaa ttcaactggc gtggttttac aacgtcgtga ctgggaaaac 360
 308 cctggcggtta cccaacttaa tcgccttgca gcacatcccc ctttcgccag ctggcgtaat 420
 310 agcgaagagg cccgcaccga tcgcccttcc caacagttgc gcagcctgaa tggcgaatgg 480
 312 gacgcgccct gtagcggcgc attaaagcgc gcgggtgtgg tggttacgcg cagcgtgacc 540
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 322 ggactcttgt tccaaactgg aacaacactc aaccctatct cggctctattc ttttgattta 840
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 332 atttccgtgt cgccttattt cccttttttg cggcattttg ccttcctgtt tttgctcacc 1140

*Xaa can only represent a single
 amino acid;
 it cannot
 represent a
 stop codon*

RAW SEQUENCE LISTING

PATENT APPLICATION: US/10/033,399

DATE: 01/18/2002

TIME: 08:04:15

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338 caatgatgag actctttaa gttctgctat gtggcgcggt attatcccgt attgacgcgc 1320
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342 cagtcacaga aaagcatctt acggatggca tgacagtaag agaattatgc agtgctgcca 1440
344 taaccatgag tgataacact gcggccaact tacttctgac aacgatcgga ggaccgaagg 1500
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350 caacaacgtt gcgcaaacta ttaactggcg aactacttac tctagcttcc cggcaacaat 1680
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356 cagcactggg gccagatggg aagccctccc gtatcgtagt tatctacacg acggggagtc 1860
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378 acaccgaact gagataccta cagcgtgagc tatgagaaag cgccacgctt cccgaaggga 2520
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386 cggccttttt acggttcttg gccttttgct ggccttttgc tcacatgttc tttcctgcgt 2760
388 tatcccttga ttctgtggat aaccgtatta ccgcctttga gtgagctgat accgctcgcc 2820
390 gcagccgaac gaccgagcgc agcagagtcag tgagcgagga agcggaagag cgccaatac 2880
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407 cagctgcagg acgtcggagg ttgcgcggcc gcagaacaaa aactgatctc agaagaggat 180
409 ctgacgcgtg ctggcggcgg ctctggtggt ggttctggtg gcggctctga ggggtggcgc 240
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413 ggttccggtg attttgatta tgaaaaaatg gcaaacgcta ataagggggc tatgaccgaa 360
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423 tgtgcgccct ttgtcttttg cgtggtgaaa ccatatgaat tttctattga ttgtgacaaa 660
425 ataaacttat tccgtggtgt ctttgcgttt cttttatatg ttgccacctt tatgtatgta 720
427 ttttctacgt ttgctaacat actgcgtaat aaggagtctt aataaggcgc gccacaattt 780

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Use of n or Naa has been detected in the Sequence Listing.
 Review the Sequence Listing to insure a corresponding
 explanation is presented in the <220> to <223> fields of
 each sequence using n or Naa.

VERIFICATION SUMMARY

PATENT APPLICATION: US/10/033,399

DATE: 01/18/2002

TIME: 08:04:16

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Output Set: N:\CRF3\01182002\J033399.raw

L:11 M:270 C: Current Application Number differs, Replaced Current Application No
L:11 M:271 C: Current Filing Date differs, Replaced Current Filing Date
L:269 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:9
L:561 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:14
L:569 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:14